

49 / ACACCCACCAIACAI CAGTGGCAATTAAGGAAGAGGTCGGAAGAGGACCTAATG 556

Db 524 ACACCACTTATCATATGCTGCAAAATTAAAGAGAGTGGAAAAAGCACTTATTG 583  
QY 557 TTGTATGCGCCATGAGATGATGGAACTGAAATGTTACTGAGAGGTGGTGGCTCTGC 616  
Db 584 TTGTATGAGGCGCATGAGATGATGGAACTGAAATGTTACTGAGAGGTGGTGGCTCTGC 643  
QY 617 TGGAAAGTGGAAAGGAAAGTGGCTGAAATGATAGCGGCGCATTTTGTGAAATACATA 676  
Db 644 TGGAAAGTGGAAAGGAAAGTGGCTGAAATGATAGCGGCGCATTTTGTGAAATACATA 703  
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QY 857 CTTCAGACTGTTTCTCACTGTACTTCGGGTAACTGGAGAAAGTTCGAACTGCTTC 916  
Db 884 CTTCAGACTGTTTCTCACTGTACTTCGGGTAACTGGAGAAAGTTCGAACTGCTTC 943  
QY 917 ACCGCTTGC----- 926  
Db 944 ACTGCTTGAAGTGGGTTTGTGATGTTCTCTGTTGTTTCCCTGGCCTCTGTGAAGAA 1003  
QY 927 ----- 926  
Db 1004 AATCACTCTAGTCCCTAAGTCACTTCTGAGCCTTGTACTGCTGGCCAGCATTTGGGC 1063  
QY 927 -----AGAGC 932  
Db 1064 CAACCGAATTTCTCCCAATCTTATCTTGGCTGGCAGAGATGTCTCAAGAGAGC 1123  
QY 933 TGATGACAGAGATGGGATTTGTTAAGTTAAATGCCAGCAATACCTGTCGAAAGCTGG 992  
Db 1124 TGATGACAGAGATGGGATTTGTTAAGTTAAATGCCAGCAATACCTGTCGAAAGCTGG 1189  
QY 993 ACTTTATCCCCAGTCTCAATTTCTGCTGGTGTCTGTATGAACAGCTTTTGTGAGAAA 1052  
Db 1184 ACTTTATCCCCAGTCTCAATTTCTGCTGGTGTCTGTATGAACAGCTTTTGTGAGAAA 1243  
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QY 1233 CTACTATATCTCAAACTTCAATTTTCTGGCCCACTCTGAGACTGATGAGAGAGATTA 1292  
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QY 1293 AGAACAGACTGAGAGATCAAGGCGCAAGAGCAAACTCAAGCTGTCTCACTGAGAGAGC 1352  
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Db 1964 GTCTGATTAAG 2023  
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QY 1893 TTTTATCTCATCTGATGAAAGTGGAGAGTGGAGAGCAATTAACAGAGAGCTTCCTT 1952  
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Db 2144 TCGGCTTTCCAGCAGCAG 2203  
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QY 2193 GCGGCGAG 2252  
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QY 2253 TTTGAAAGAGATTTAAAG 2312  
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2744 TCTTACATCTGATTTGAGATGATGATTAATCTCTCAACAAAATGGAAGGAG 2803  
2613 ATGCTAGAAATCCCCCTAGAGAGGAGAAATTTATTCATGAAATAGCATCCCT 2672  
2804 ATGCTAGAAATCCCCCTAGAGAGGAGAAATTTATTCATGAAATAGCATCCCT 2863  
2673 TGTCTTAAATGAGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 2732  
2864 TGTCTTAAATGAGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 2922  
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3223 ATCTGTAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 3182  
3093 CCAATTCAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 3152  
3283 CCAATTCAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 3242  
3153 CTTCTCAGCTTATGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 3212  
3243 CTTCTCAGCTTATGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 3302  
3213 TGGCAAGTACAGATTTAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 3272  
3403 TGGCAAGTACAGATTTAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 3362  
3273 AATCAAGATTTGTTAAATGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 3332  
3463 AATCAAGATTTGTTAAATGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 3422

RESULT 2  
US-09-816-494-3  
Sequence 3, Application US/09816494  
Patent No. 6664089  
GENERAL INFORMATION:  
APPLICANT: Weiers, Rachel A.  
TITLE OF INVENTION: 38692 AND 21117, NOVEL DUAL SPECIFICITY  
FILE REFERENCE: 10448-030002  
CURRENT APPLICATION NUMBER: US/09/816,494  
PRIORITY FILING DATE: 2001-03-23  
PRIOR APPLICATION NUMBER: US 60/191,858  
NUMBER OF SEQ ID NOS: 10  
SOFTWARE: FASTSEQ for Windows Version 4.0  
SEQ ID NO 3  
LENGTH: 1998  
TYPE: DNA  
ORGANISM: Homo sapiens  
US-09-816-494-3

Query Match 49.8%; Score 1660; DB 4; Length 1998;  
Best Local Similarity 91.8%; Pred. No. 0;  
Matches 1884; Conservative 0; Mismatches 0; Indels 164; Gaps 1;  
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1 ATGGCCCTTATGAGATTTGAAATCTCAATTTGTTACTGAGAGGTTGGCTTCTGAGAA 60  
622 ATGGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 681  
61 ATGGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 120  
682 CACATTTTGAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 741  
121 CACATTTTGAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 180  
742 GACAAAGTTTAAATTAAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 801  
181 GACAAAGTTTAAATTAAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 240  
802 TGCAGTGAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 861  
241 TGCAGTGAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 300  
862 GACTGTTTCTCATCTGATCTTCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 921  
301 GACTGTTTCTCATCTGATCTTCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 360  
922 CTGTC----- 926  
361 CTGTCAGTGGGTTTCTGATCTTCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 420  
927 ----- 926  
421 ACTGATGCTTACCTGATTTCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 480  
927 ----- 926  
481 GCAATTCCTCCAAATTTTATCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 540  
938 CAGCAAGTGAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 997  
541 CAGCAAGTGAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 600  
998 ATCCCGAGGCTCATTTCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1057  
601 ATCCCGAGGCTCATTTCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 660  
1058 CCGTGTGAGCAAAATCAGTAGATTTCAATGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1117  
661 CCGTGTGAGCAAAATCAGTAGATTTCAATGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 720  
1118 CTAGTGACATGTTTACCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1177  
721 CTAGTGACATGTTTACCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 780  
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1238 ATATCTCAAAATCTCAATTTCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1297  
841 ATATCTCAAAATCTCAATTTCTGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 900  
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901 CAGACTGAGCATCAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 960  
1358 GAACTTCTCTGCTGCTCAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1417  
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Db      592 CCGACCTTATCCCGAGCTCATCTTCTCGCTGCTCCGTAATGACACCTTTTGAG 651
Qy      70  Lys11leuProtrPleuAspLysSerValAspPhe11eGluValAlaValAspAen 89
Db      652 AAAATTTGCCGTGGTGGACAAATCAGTATGATTTCTATGAGAAAGCAAAAGCTCCAAAT 711
Qy      90  GlyCyvalLeuValHisCyaleuAlaGly11SerArgSerAlaTrilleAla11eAla 109
Db      712 GGAATGTTCTAGTGAAGCTTTAGCTGGGATCTCCCGCTCCGCAACCTCCATCTGAGC 771
Qy      110 Tyrl1eMetLysArgMetLysPheSerLeuAspGluAla17YrhArgPheVal1yGluLys 129
Db      772 TACATCATGAGAGATGACATGATCTTTAGTGAAGCTTACAGATTGTGAAGAAAAA 831
Qy      130 ArgProThr11eSerProAspPheAsnPheLeuGlyGlnLeuLeuAspTYR31uLysLys 149
Db      832 AGACCTACTATCTCCAACTCAATTTCTGGGCCAACTCTGGACTATGAGAAAGAAAG 891
Qy      150 11eLysAsnGlnTh1GlyAlaSerGlyProLysSerLysLeuLysLeuLys11eLysGlu 169
Db      892 ATTAAAGAACCAAGCTGAGCATCAAGGCCAAAGCAAACTCAAGCTGCTGCACTGAG 951
Qy      170 LysProAsnGluProValProAlaValSerGluGlyGlnLysSerGluThrProLeu 189
Db      952 AAGCCAAATTAACCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1011
Qy      190 SerProProCysAlaAspSerAlaThrSerGluAla11eGlyGlnArgProVal11ePro 209
Db      1012 AGTCAACCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1071
Qy      210 AlaSerValProSerValProSerValGlnProSerLeuLeuGluAspSerProLeuVal 229
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Qy      230 G1AlaLeuSerGlyLeuHisLeuSerAlaAspArgLeuGluAspSerLeuLysLys 249
Db      1132 CAGCGCTCGAGGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1191
Qy      250 ArgSerPheSerLeuAspLysLysSerValSerTYRserAlaSerMetAla11eSerLeu 269
Db      1192 CGTTCCTCTCTGATATCAAAATCAGTTTCTATTCAGCGCAGAGGAGGAGCATCTCTTA 1251
Qy      270 HisGlyPheSerSerSerGluAspAlaLeuGluTYR1YrhYrhProSerThrLeuAsp 289
Db      1252 CAGGCTTCTCTCTCAAGAGAGAGCTTGAATACCAAACTTCACTGATCTGAT 1311
Qy      290 G1YrhAspLysLeuGlnPheSerProValGlnGluLeuSerGluGln11eThrProGlu 309
Db      1312 GGAACCAACAGCTATGCAATCTTCTCCCTGCTGCAAGACTATCGAGCAAGACTCCGAA 1371
Qy      310 ThrSerProAspLysGluGluAlaSerT1eProLysLysLeuGlnThrAlaArgProSer 329
Db      1372 ACCAGTCTGATTAAGAGAGAGCAGCATCCCAAAAGACTGAGACCGCCAGGCTCTTA 1431
Qy      330 AspSerGlnSerLysArgLeuHisSerValArgTYRserSerSerGlyTYRAlaGlnArg 349
Db      1432 GACAGCCAGAGCAGAGGATGATTCATTCGGTCAAGACAGAGAGAGAGAGCCCGCAGAG 1491
Qy      350 SerLeuLeuSerProLeuHisArgSerGlySerValGluAspAsnTYRHisArgSerPhe 369
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Qy      370 LeuPheGlyLeuSerThrSerGlnGlnHisLeuThrLysSerAlaGlyLeuGlyLeuLys 389
Db      1552 CTTTCGGGCTTTCACAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1611
Qy      390 GlyTYRHisArgSerPheLeuAlaProGlnThrSerThrProSerLeuThrSerSerTYR 409
Db      1612 GCGGAGACTCGAATATCTGCGCCCAAGACCTTCAACCTTCCCTGAGACAGACACTGAG 1671
Qy      410 TyrlPheAlaThrGluSerSerHisPheTYRserAlaSerAla11eTYRGLYGLYSerAla 429

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Db      1672 TACTTGCAACAGACTCTCACTTACTCTGCTCAGCCACTATAGAGAGAGACTGCC 1731
Qy      430 SerTYRserAlaTYRserCysSerGlnLeuProThrCysGlyAspGlnValTYRserVal 449
Db      1732 AGTTACTGCTGCTTACAGCTGAGCCAGCTGCCACTTCCGAGAGACCAAGTCTTCTG 1791
Qy      450 ArgArgArgGlnLysProSerAspArgAlaAspSerArgTYRserTYRHisArgGluSer 469
Db      1792 CGCAGCGCCAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1851
Qy      470 ProPheGluLysGlnPheLysArgArgSerCysGlnMetGluPheGlyGluSerT1eMet 489
Db      1852 CCGTTGAAAAGAGCTTTAAACGAGAGCTGCAAAATGAAATTTGAGAGAGAGAGAGAG 1911
Qy      490 SerGluAspArgSerArgGluGluLysVal1GlySerGlnSerSerPheSerGly 509
Db      1912 TCAGAGAACAGGTCACGGGAGAGAGCTGAGGAGAAAGTGGCACTGATCTTCTGAGC 1971
Qy      510 SerMetGluLys11eGluValSer 517
Db      1972 AGCATGGAATCATGAGCTTCC 1995

RESULT 2
US-09-816-494-1
Sequence 1, Application US/09816494
Patent No. 6664089
GENERAL INFORMATION:
APPLICANT: Meyers, Rachel A.
TITLE OF INVENTION: 38692 AND 21117, NOVEL DUAL SPECIFICITY
FILE REFERENCE: 10448-030002
CURRENT APPLICATION NUMBER: US/09/816,494
PRIOR FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: US 60/191,858
PRIORITY FILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO: 1
LENGTH: 3544
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (589) ... (2563)
US-09-816-494-1

Alignment Scores:
Pred. No.: 0 Length: 3544
Score: 488.00 Matches: 488
Percent Similarity: 100.00% Mismatches: 0
Best Local Similarity: 100.00% Indels: 0
Query Match: 94.35% Gaps: 0
DB: 4

US-09-964-277-21 (1-517) x US-09-816-494-1 (1-3544)
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Db      1120 GAGCTATGACACCAAGATGAGATGCTTGGTAAATGCCACCAATATCTCTCCAAAG 1179
Qy      50  ProAspPhe11eProGluSerHisPheLeuArgValProValAsnAspSerPheCysGlu 69
Db      1180 CCGATCTTATCCCGAGATCCATTCCTGCGTGGCGCTCGATGACACTTTTGAG 1239
Qy      70  Lys11leuProtrPleuAspLysSerValAspPhe11eGluValAlaValAspAen 89
Db      1240 AAAATTTGCCGTGGTGGACAAATCAGTATGATTTCTATGAGAAAGCAAAAGCTCCAAAT 1299
Qy      90  GlyCyvalLeuValHisCyaleuAlaGly11SerArgSerAlaTrilleAla11eAla 109
Db      1300 GGAATGTTCTAGTGAAGCTTTAGCTGGGATCTCCCGCTCCGCAACCTCCATCTGAGC 1359
Qy      110 Tyrl1eMetLysArgMetLysPheSerLeuAspGluAlaTYRArgPheValLysGluLys 129

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Db      1420 AGACCTACTATATCTCCAACTTCAATTTCTGGGCGCAACTCTGACATATGAGAAAG 1479
Qy      150  IleValAsnGlnThrGlyAlaSerGlyProLysSerLeuLeuLeuLeuLeuGln 169
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Qy      170  LysProAsnGluProValProAlaValSerGluGlyGlnLeuYsSerGluThrProLeu 189
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Qy      190  SerProProCysAlaAspSerAlaThrSerGluAlaAlaGlyGlnArgProValHisPro 209
Db      1600 AGTCCACCTGTGCGGACTCTGCTACTCTGAGGAGCAGGACGAAAGCCCGTGCATCC 1659
Qy      210  AlaSerValProSerValProSerValGlnProSerLeuLeuGlnLysSerProLeuVal 229
Db      1660 GCCAGGCTGCCAGGCTGCCAGGCTGCCAGGCTGCCAGGCTGCCAGGCTGCCAGG 1719
Qy      230  GlnAlaLeuSerGlyLeuHisLeuSerAlaAspArgLeuGlnLysSerAlaLeuYs 249
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Qy      250  ArgSerPheSerLeuAspIleYsSerValSerYsSerAlaSerMetAlaLeuSerLeu 269
Db      1780 CGTCTCTCTCTCTGATATCAATCATATCATATTCATATTCAGCAGCATGCGACATCTTA 1839
Qy      270  HisGlyPheSerSerSerGluAspAlaLeuGlnYsYsYsYsYsYsYsYsYsYsYs 289
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Qy      290  GlyThrAlaLeuYsLeuYsGlnPheSerProValGlnGlnLeuSerGluGlnThrProGln 309
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Qy      310  ThrSerProAspLysGluGlnAlaSerIleProYsLysLeuGlnThrAlaArgProSer 329
Db      1960 ACCAGCTCTGATTAAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 2019
Qy      330  AspSerGlnSerLeuYsArgLeuHisSerValArgThrSerSerSerGlyThrAlaArg 349
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Db      2140 CTTTTCGCGCTTTCACACAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAG 2199
Qy      390  GlyTPHAspAspIleLeuAlaProGlnThrSerThrProSerLeuHisSerSerThr 409
Db      2200 GGTGGGAGCTCGATATCTTGGGCGCGCGAGCCTTACCTTCCCTGACCGAGGAGG 2259
Qy      410  TyrPheAlaThrGlnLysSerThrHisPheYsSerAlaSerAlaIleYsGlyYsSerAla 429
Db      2260 TATTTGCGACAGAGTCTCACTGCTCACTGCTCACTGCTCACTGCTCACTGCTCACT 2319
Qy      430  SerYsSerAlaYsSerYsSerGlnLeuProThrCysGlyAspGlnValYsSerVal 449
Db      2320 ACTTACTCTGCTCACTGCTCACTGCTCACTGCTCACTGCTCACTGCTCACTGCTG 2379
Qy      450  ArgGAGArgGlnLysProSerAspArgAlaAspSerArgYsSerThrIleGluGlnYs 469
Db      2380 CCGAGGCGGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 2439
Qy      470  ProPheGlnYsGlnPheYsArgArgSerCysGlnMetGlnPheGlyLeuSerIleMet 489

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Db      2440 CCCTTGAAGAGCAGTTTAAACGAGAGAGCTGCCAATGGAATTGGAGAGAGCATCAG 2499
Qy      490  SerGluAsnArgSerArgGlnGlnLeuGlnYsValGlnYsSerGlnSerSerPheSerGly 509
Db      2500 TCGAGGAGCAGTCAACGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 2559
Qy      510  SerMetGlnIleIleGluValSer 517
Db      2560 AGCATGAGAAATCATTTGAGGCTTCC 2583

RESULT 3
US-09-920-668-3
Sequence 3, Application US/09920668
Patent No. 6482644
GENERAL INFORMATION:
APPLICANT: Lex M. Cowart
TITLE OF INVENTION: ANTISENSE MODULATION OF DUAL SPECIFIC PHOSPHATASE 8 EXPRESSION
FILE REFERENCE: RTS-0246
CURRENT APPLICATION NUMBER: US/09/920,668
NUMBER OF SEQ ID NOS: 49
SEQ ID NO 3
LENGTH: 2377
TYPE: DNA
ORGANISM: Homo sapiens
FEATURES:
NAME/KEY: CDS
LOCATION: (135) ... (2012)
US-09-920-668-3

Alignment Scores:
Pred. No: 2,786-10 Length: 2377
Score: 20.00 Matches: 20
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 3.87% Indels: 0
DB: 4 Gaps: 0

US-09-964-277-21 (1-517) x US-09-920-668-3 (1-2377)

Qy      94  ValHisCysLeuAlaGlyIleSerArgSerAlaThrIleAlaIleAlaYsIleMetYs 113
Db      864  GTCCACTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 923

RESULT 4
US-09-016-434-1135
Sequence 1135, Application US/09016434
Patent No. 6500938
GENERAL INFORMATION:
APPLICANT: Janice Au-Young
APPLICANT: Jeffrey J. Sellhammer
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF SIGNALING
TITLE OF INVENTION: PATHWAY GENE EXPRESSION
NUMBER OF SEQUENCES: 1490
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
STREET: 3174 FORSTER DRIVE
CITY: PALO ALTO
STATE: CALIFORNIA
COUNTRY: USA
ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/016,434
FILING DATE: HERewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:

```